

# Small cell lung cancer

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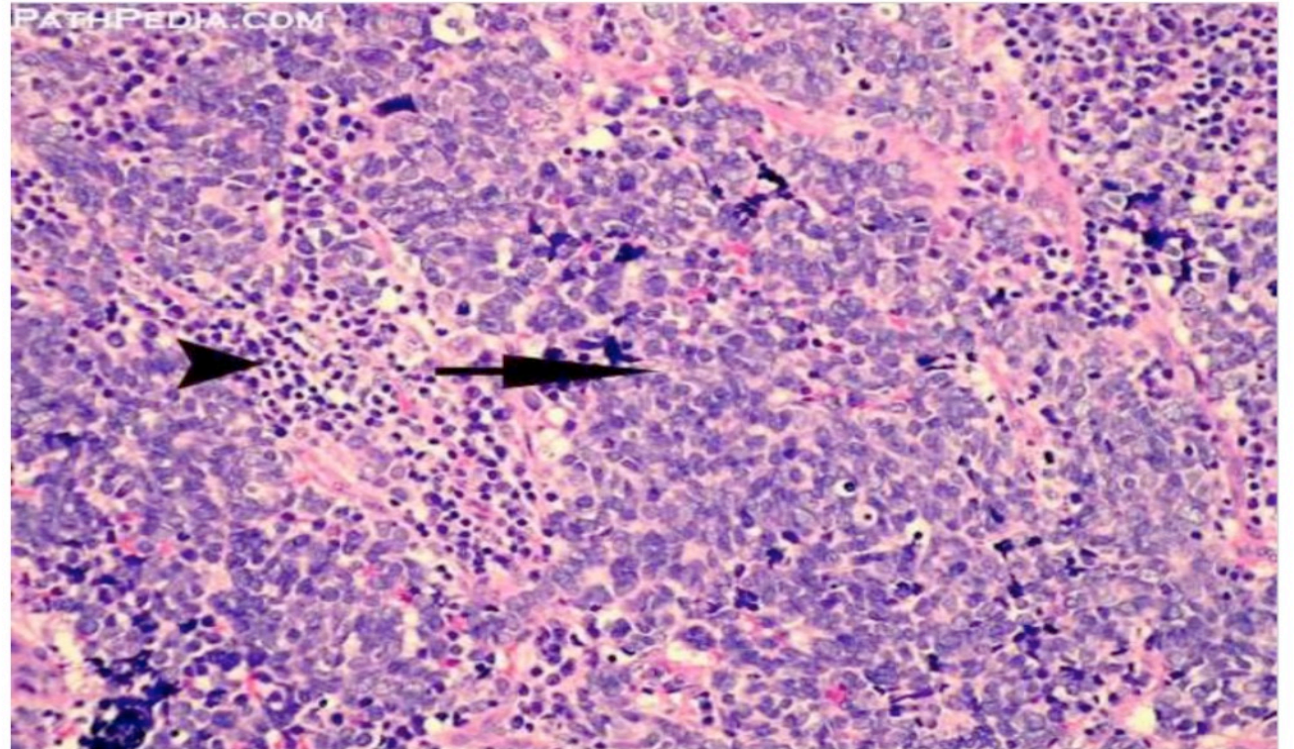
# Outline

- Small cell lung cancer 101
- Genetic abnormalities of small cell lung cancer
- SCLC as a resistance mechanism to EGFR TKI in lung adenocarcinoma
- Examples of translational medicine: Story of Rova-T
- Examples of translational medicine: Immune checkpoint inhibitors
- Other promising agents under clinical development
- Extrapulmonary small cell carcinoma

# SCLC morphology

## Morphology of SCLC

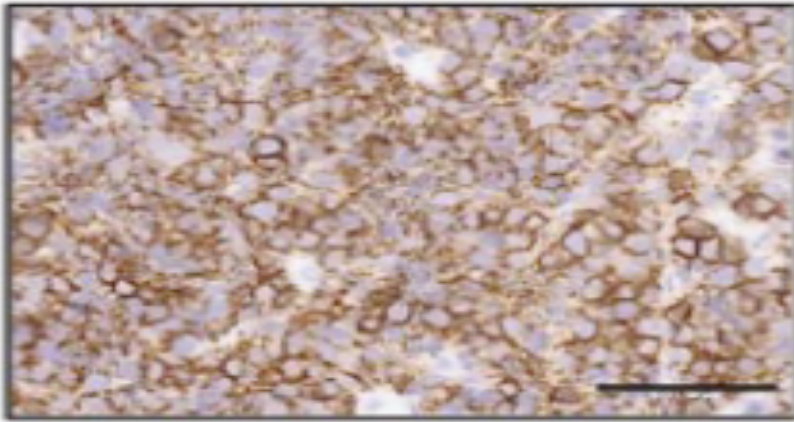
Small cell lung cancer (SCLC) is also known as oat cell carcinoma. Its morphology resembles oat grains and appears as small oval cells with scanty cytoplasm.



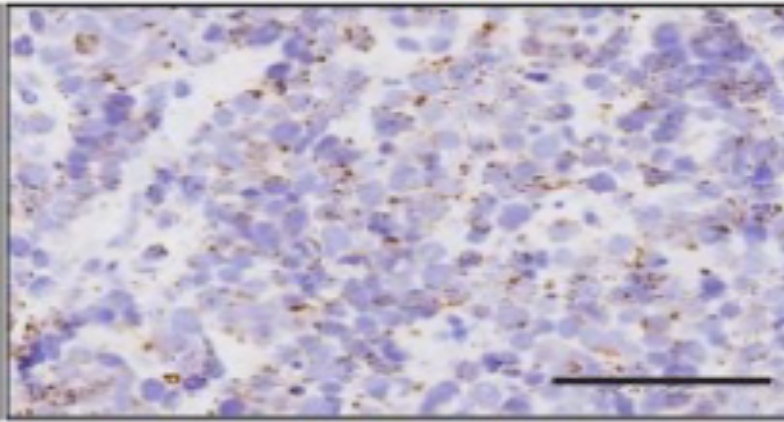
# IHC staining

## IHC staining to diagnose SCLC

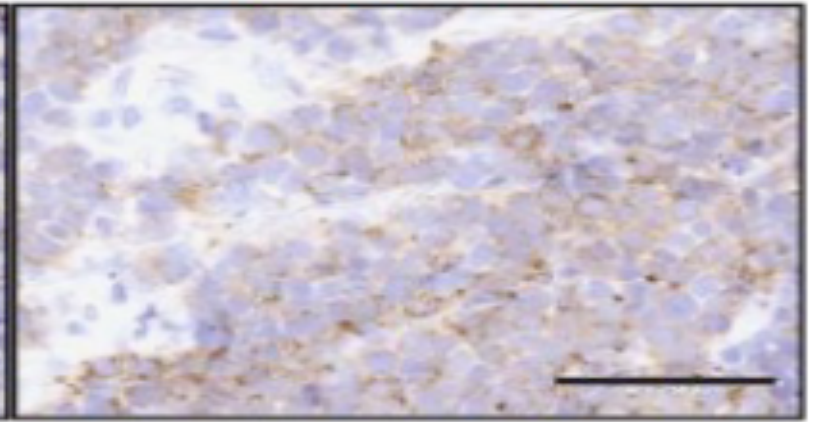
CD56



Chromogranin



Synaptophysin





# SCLC

## SCLC

- SCLC accounts for 10% to 15% of all lung cancer cases, and is closely linked to the intensity and duration of tobacco smoking.
- Compared to NSCLC, SCLC tends to disseminate earlier in the course of its natural history and displays a more aggressive clinical behavior.
- SCLC is also commonly associated with paraneoplastic endocrinopathies (e.g., Cushing syndrome).

# SCLC is considered as a recalcitrant cancer

- Recalcitrant Cancer Research Act of 2012.
- Recalcitrant cancer:
  - Have a 5-year relative survival rate of less than 20%
  - Estimated to cause the death of at least 30,000 individuals in the United States per year.
- NCI identified four major obstacles to progress in 2014:
  - Continuing risk of developing the disease that remains for decades after smoking cessation.
  - Most patients have widely metastatic tumors at the time of diagnosis.
  - Rapid development of resistance to chemotherapy in more than 95% of SCLC patients.
  - Lack of tumor tissue for clinical, molecular , and cell biological studies.

SCLC:

<7%

~30,000  
deaths/yr

# Systemic therapy of SCLC

- It was learned quite early in the 1970s that combination therapy produces superior survival compared with single-agent treatment based on several randomized trials.
- First-line therapy: platinum + etoposide
- Second-line therapy: Topotecan
- Third line therapy: Nivolumab

# EPSCC

- Genetic abnormalities of SCLC

## Extrapulmonary small cell carcinoma (EPSCC)

**Table 1**  
Frequency of EPSCC per site of origin.

	Percentage of SCC/total per site of origin	Estimated number of patients in US per year*
Pulmonary	15–20%	32,250–43,000
Oesophagus	0.8–2.4%	130–395
Larynx	0.5–1%	60–120
Bladder	0.3–1.0%	200–680
Cervix	±1%	±110
Prostate	±2%	±250
Unknown primary	7–30% of all EPSCC	70–300

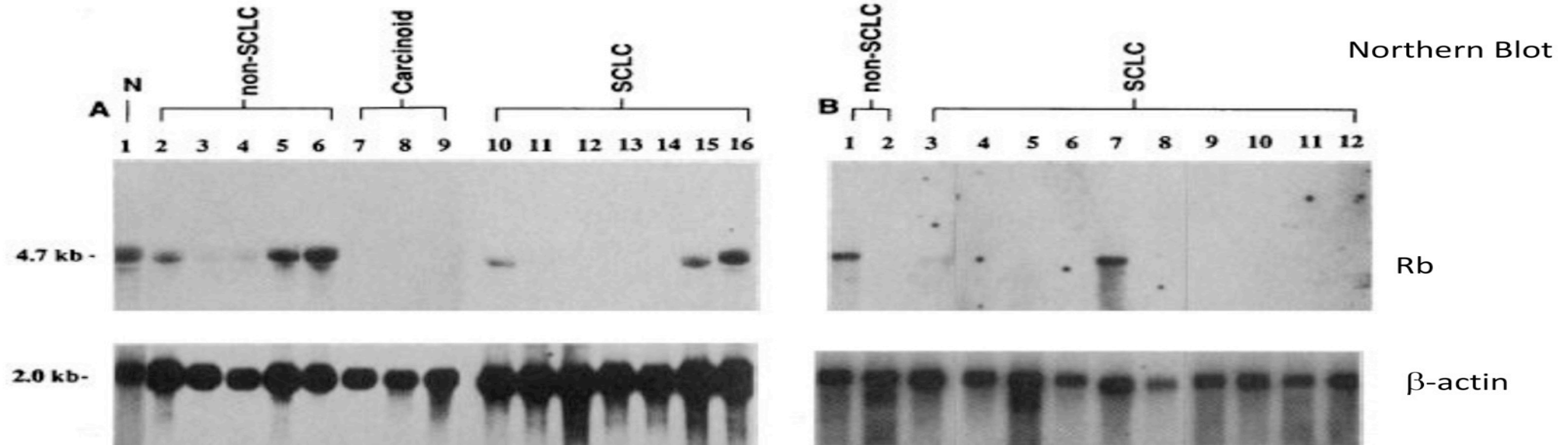
SCC denotes small cell carcinoma; EPSCC denotes extrapulmonary small cell carcinoma.

\* <http://www.cancer.gov/cancertopics/pdq>.



# RB loss

## Genetic abnormalities of SCLC— Loss of Rb gene



# TP53 inactivation

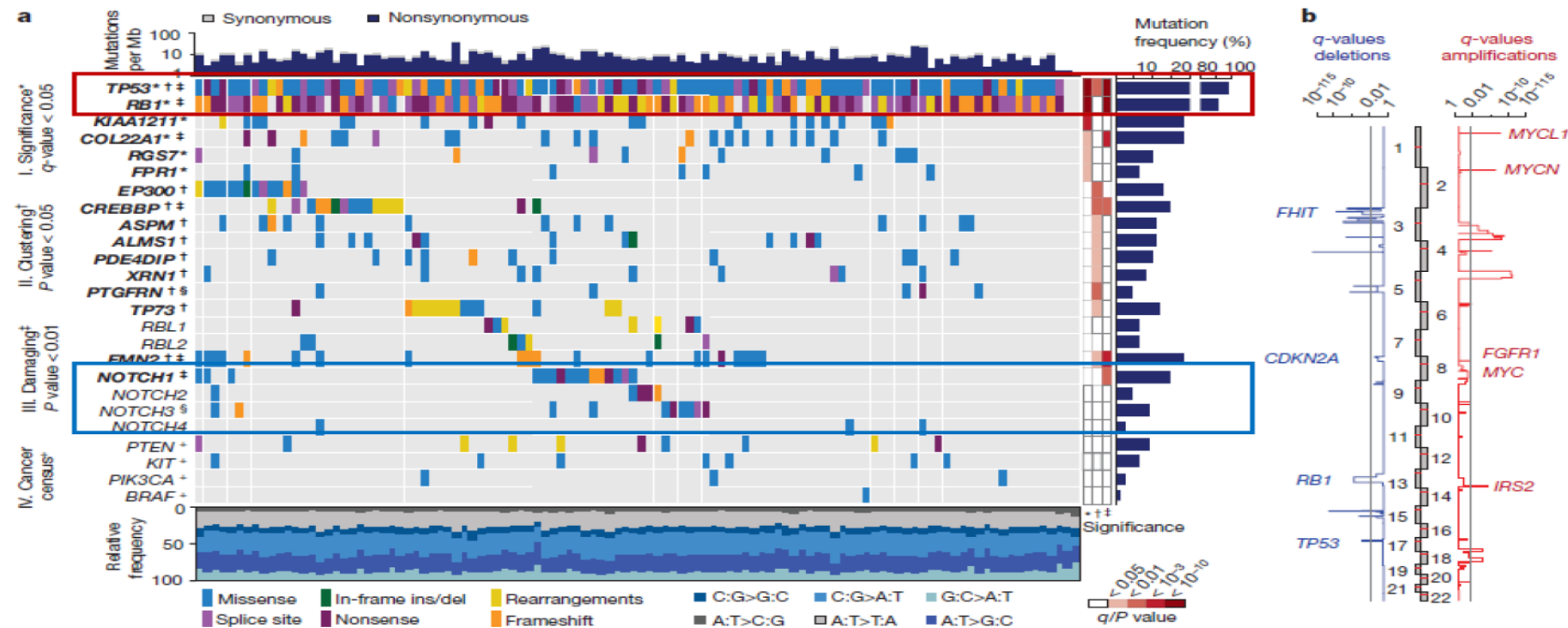
## Genetic abnormalities of SCLC – Inactivation of TP53 gene

**Table 1.** Abnormalities of p53 in lung cancer lines. Terms and symbols for mRNA levels are as follows: +, easily detectable p53 transcripts comparable to levels found in normal lung; reduced or trace, greatly reduced amount of transcript compared to normal lung; undetectable, undetectable by both Northern blot analysis and the RNase protection assay. Full designation of the cell lines includes the prefix “NCI”. All but H60, H69, H82, H187, H345, H378, and H510 were established from patients before treatment.

Type of mutations	mRNA level	Tumor cell type	Cell line
Homozygous deletion	Undetectable	Bronchioloalveolar	H358
Homozygous deletion with truncated mRNA	Reduced	Extrapulmonary small cell	H660
DNA rearrangement	Undetectable	Adenocarcinoma	H969
Abnormal size mRNA	+	Small cell	H526
	+	Adenocarcinoma	H676
	+	Adenosquamous	H647
	Trace	Small cell	H82
Point or small mutation	+	Small cell	H1436, H1450
	+	Pulmonary carcinoid	H727
	+	Adenocarcinoma	H23
	+	Bronchioloalveolar	H820
	+	Adenosquamous	H125
	+	Large cell	H661
	Reduced	Small cell	H889, H1092
	Reduced	Adenocarcinoma	H920
None detected	Trace	Small cell	H60, H69, H209, N417
	Reduced	Squamous	H520
None detected	+	Small cell	H187, H345, H378
	+	Extrapulmonary small cell	H510
	+	Adenosquamous	H596
	+	Squamous	H226
	+	Large cell	H460, H1385

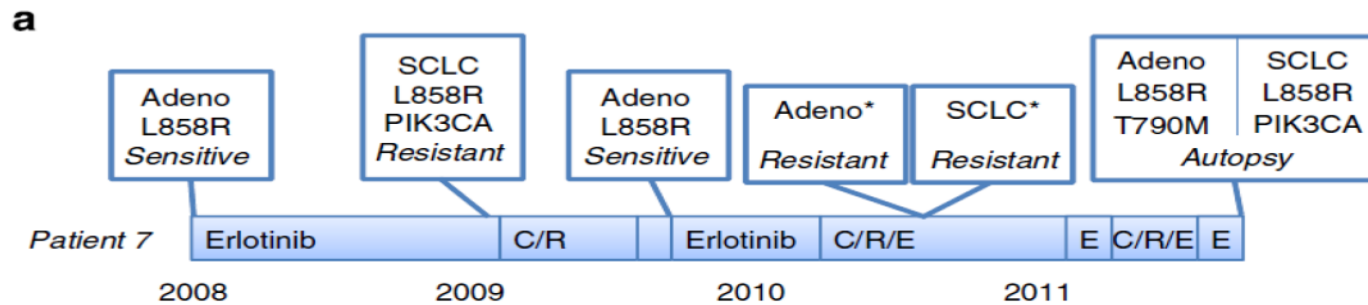
# Genetic abnormalities

## Genetic abnormalities of SCLC: WES Analysis



# SCLC conversion

SCLC conversion as a resistance mechanism to EGFR TKI in lung adenocarcinoma: Loss of TP53 and Rb genes



**b**

Sample	Normal liver	Diaphragm tumour	Lung tumour	Liver tumour
Histological features	Normal tissue	Adenocarcinoma	SCLC	SCLC
Number of reads	179,298,190	350,864,233	388,189,232	318,482,313
Average coverage	146	287	319	253
Primary <i>EGFR</i> mutation	WT	L858R	L858R	L858R
Secondary <i>EGFR</i> mutation	WT	T790M	WT	WT
<i>PIK3CA</i> status	WT	WT	E545K	E545K
<i>TP53</i> status	WT	WT/Δ154–163	–/Δ154–163	–/Δ154–163

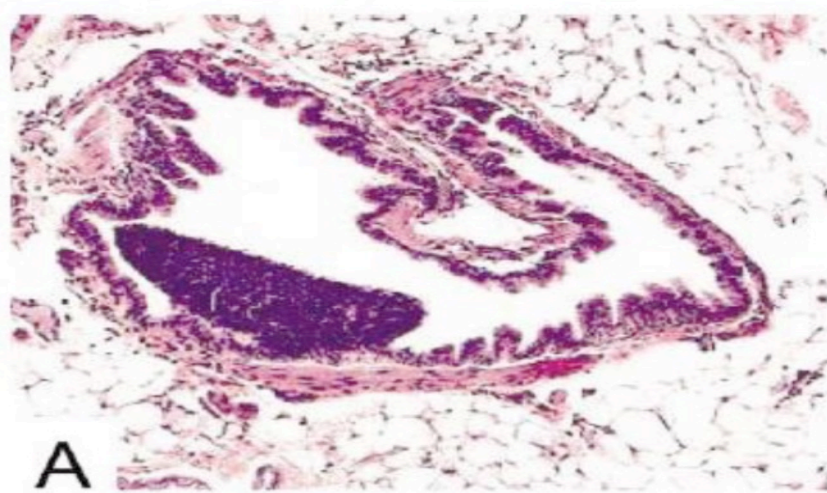
**Table 1 | RB status of TKI-resistant patients.**

Patient	Cancer type	Resistance	Histology	RB status	Detection method
1	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	NE	Neg	IHC/genetic
	Lung	Post	NE	Neg	IHC/genetic
2	Lung	Pre	Adeno	Pos	IHC
	Lung	Pre	Adeno	Neg	IHC
	Lung	Post	NE	Neg	IHC
3	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	NE	Neg	IHC
4	Lung	Post	NE	Neg	IHC
5	Lung	Post	NE	Neg	IHC
6	Lung	Pre	Adeno	Neg	IHC
	Lung	Post	NE	Neg	IHC/genetic*
7	Lung	Post	Adeno	Pos	IHC/genetic
	Lung	Post	NE	Neg	IHC/genetic
	Lung	Post	NE	Neg	Genetic
8	Lung	Post	Adeno	Pos	IHC
	Lung	Post	NE	Neg	IHC
9	Lung	Post	NE	Neg	IHC
10	Lung	Post	Adeno	Neg	IHC
11	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	Adeno	Pos	IHC
12	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	Adeno	Pos	IHC
13	Lung	Post	Adeno	Pos	IHC
14	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	Adeno	Pos	IHC
15	Lung	Post	Adeno	Pos	IHC
16	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	Adeno	Pos	IHC
17	Lung	Pre	Adeno	Pos	IHC
	Lung	Post	Adeno	Pos	IHC
18	Lung	Post	Adeno	Pos	IHC
19†	Lung	Intrinsic	NE	Neg	IHC

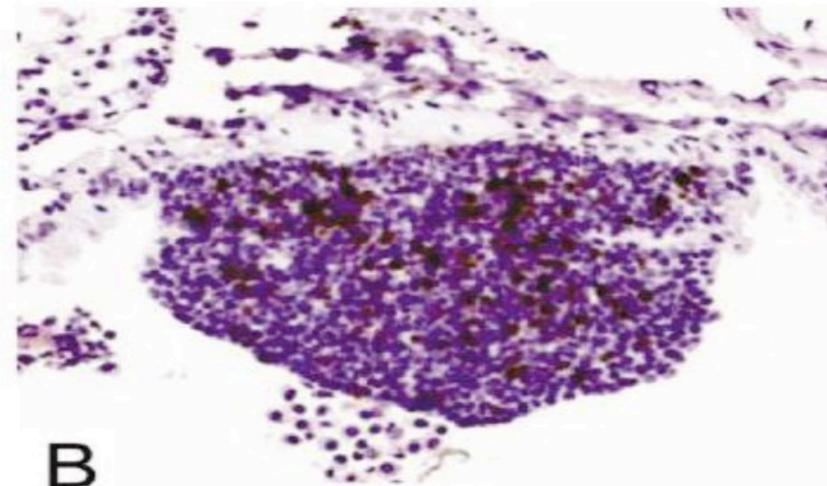


# TP53 and Rb inactivation

Conditional inactivation of Trp53 and Rb1 led to SCLC in mouse model



A  
Hyperplastic focus in the airway  
(H&E staining)



B  
Anti-BrdU staining

SCLC became detectable within 196-350 days in the mouse model with conditional KO of TP53 and Rb1.

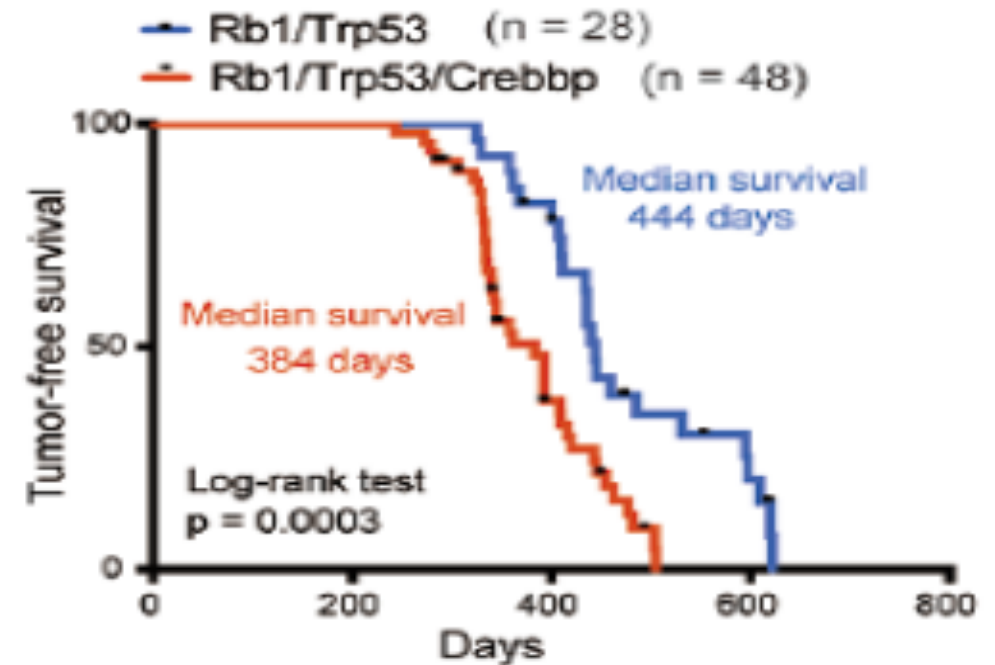
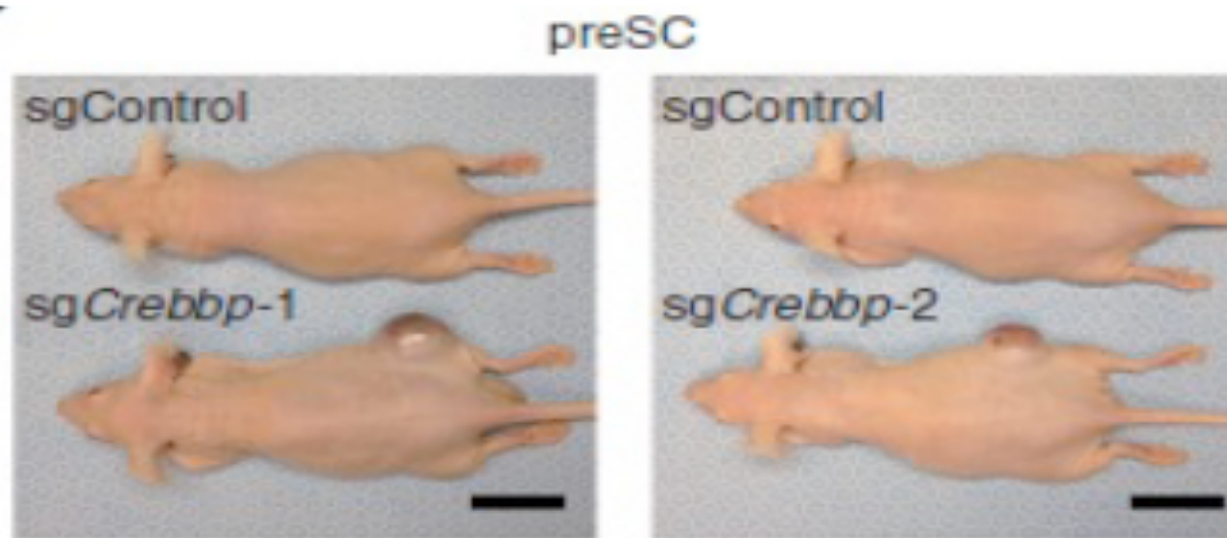
# Genomic abnormalities

## Genomic abnormalities of SCLC

1. Inactivation of Rb and TP53
2. Inactivation of Epigenetic genes EP300 and CREBBP
3. Inactivation of Notch signaling

# Inactivation of Crebbp

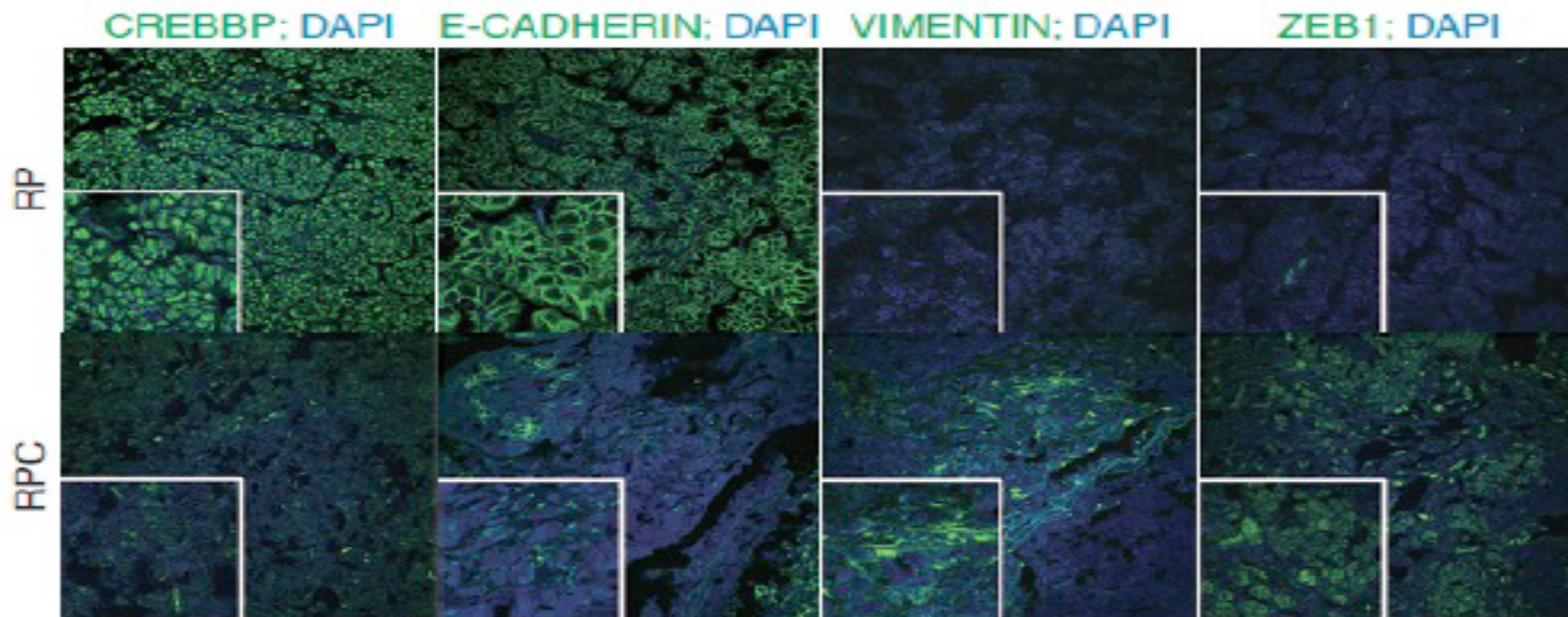
Inactivation of Crebbp accelerated development of SCLC





# E-Cadherin

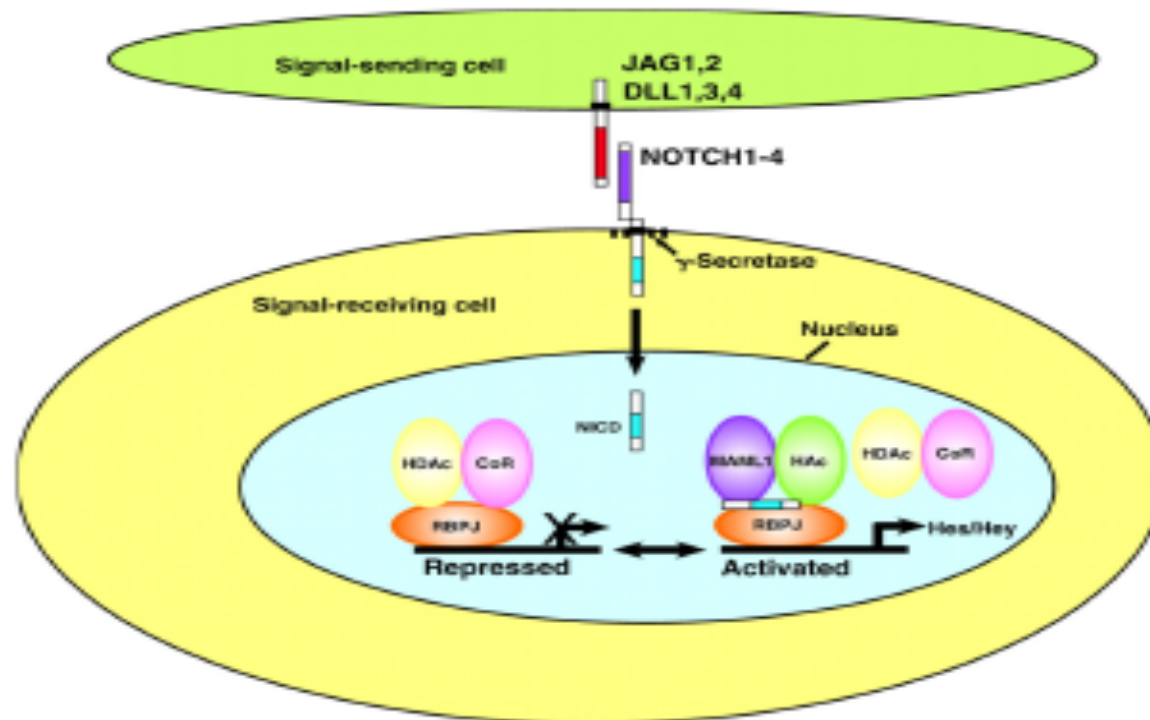
## Inactivation of Crebbp Reduced Expression of E-cadherin





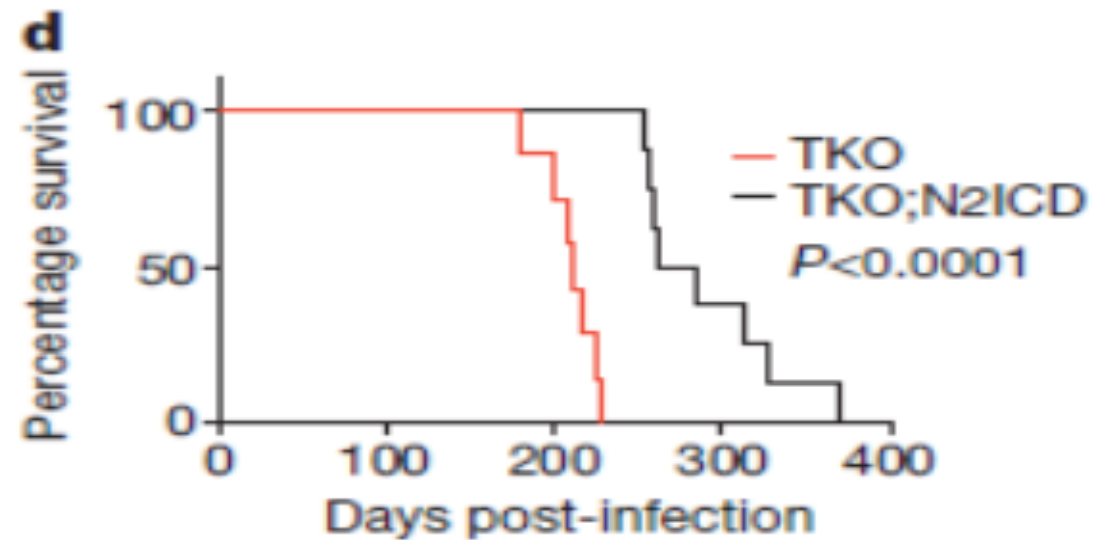
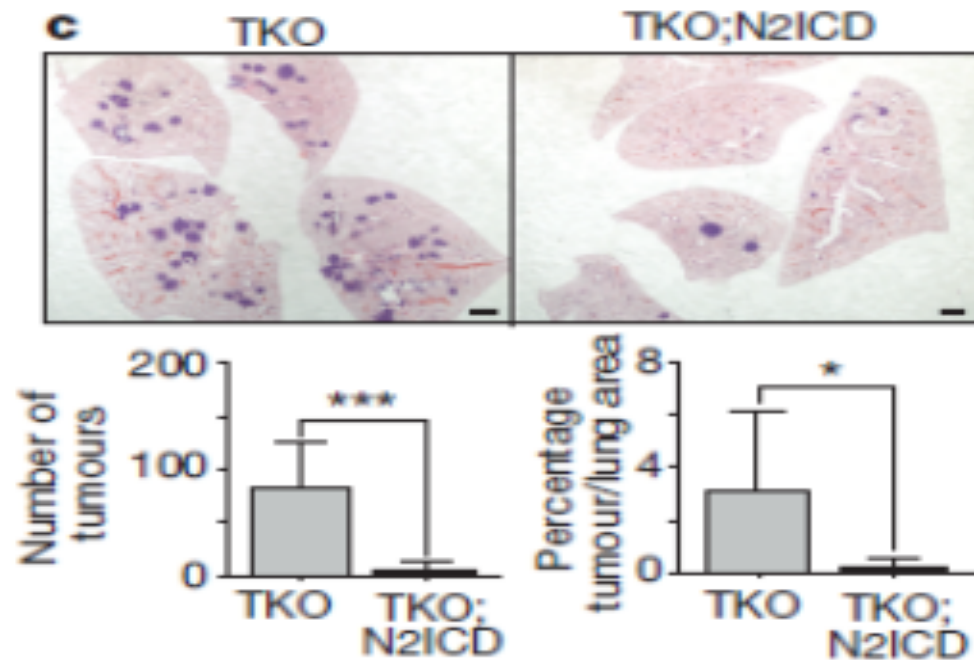
# Notch signaling pathway

## Notch Signaling Pathway



# Notch signaling

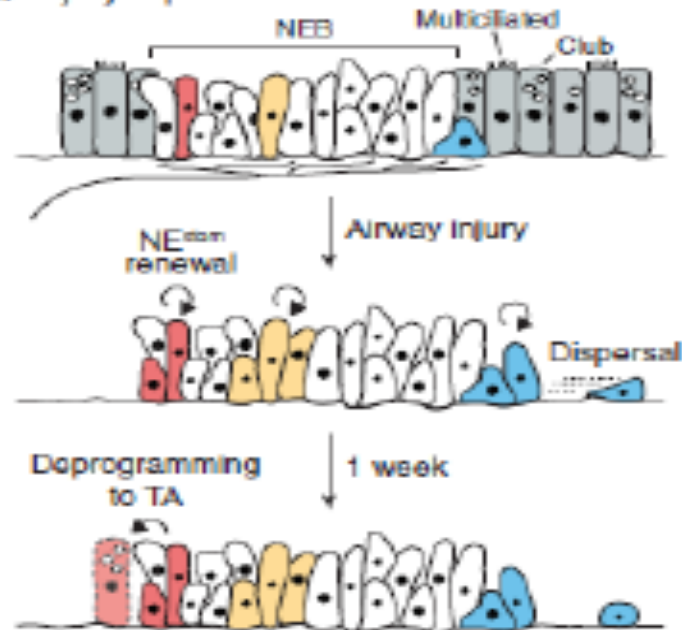
Forced activation of Notch signaling decreased SCLC growth in a transgenic mouse model



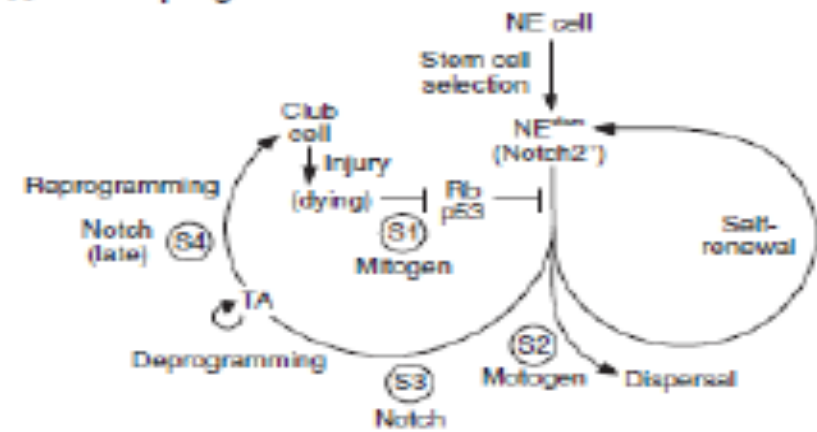
# Mouse model using NE stem cells

A model: SCLC is developed from NE stem cells

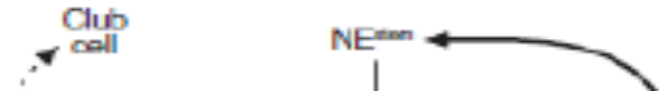
G Injury repair



H NE<sup>stem</sup> program



I Small cell lung cancer

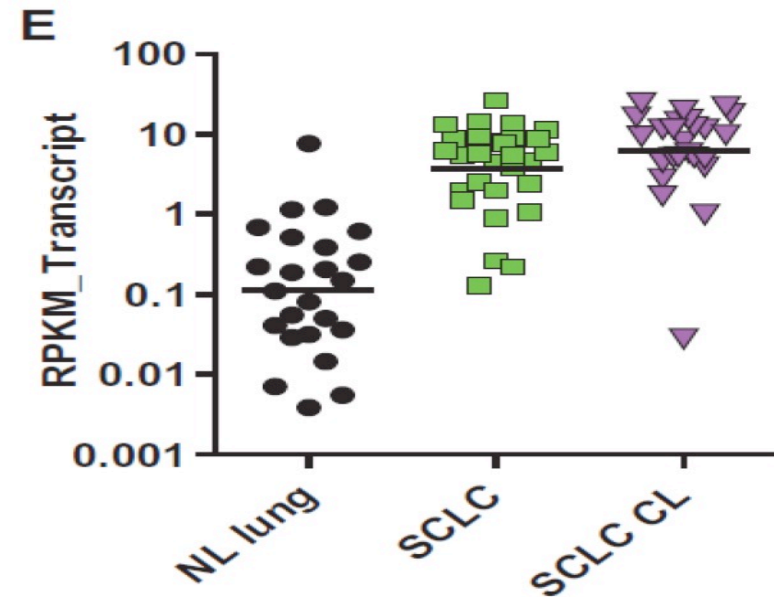
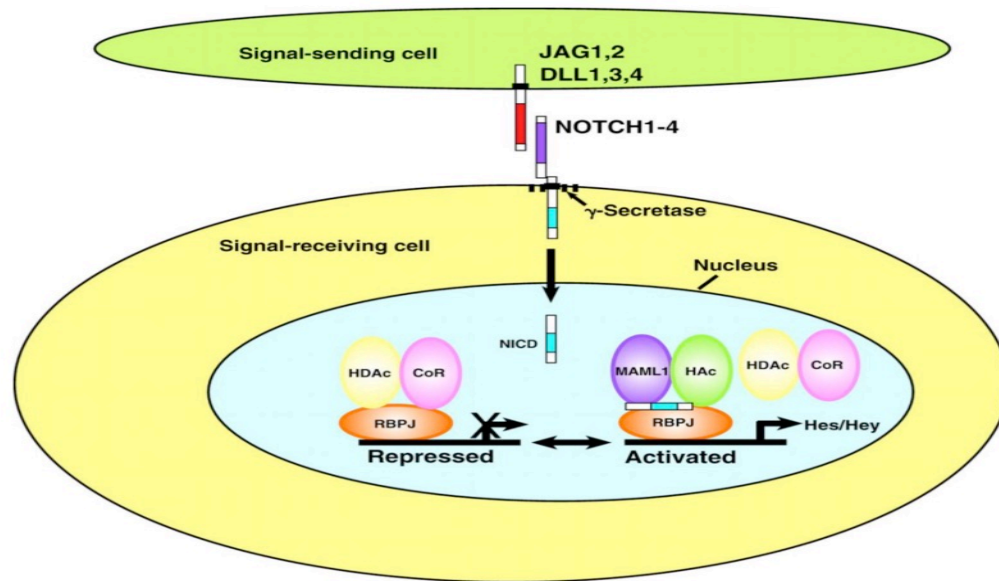


# Examples of Translational medicine: Story of Rova-T



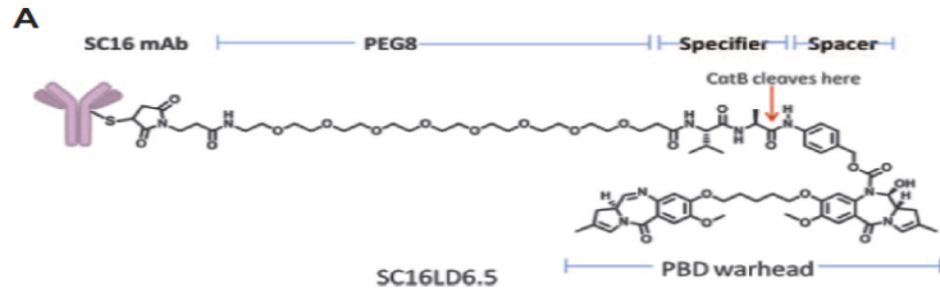
# DLL3 overexpression

## Overexpression of DLL3 in SCLC

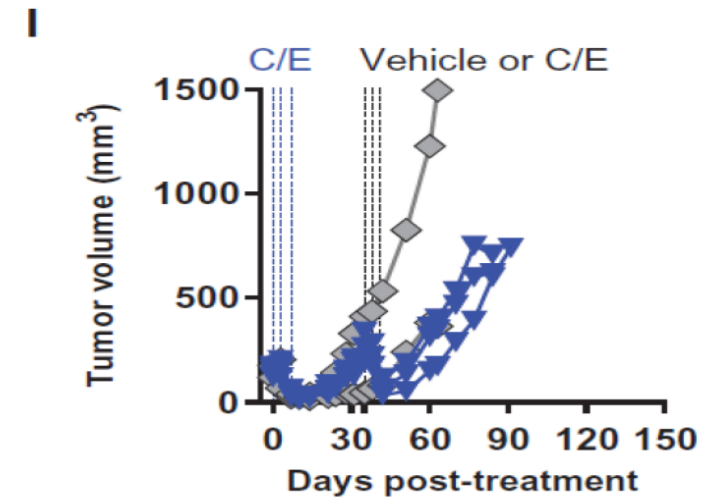
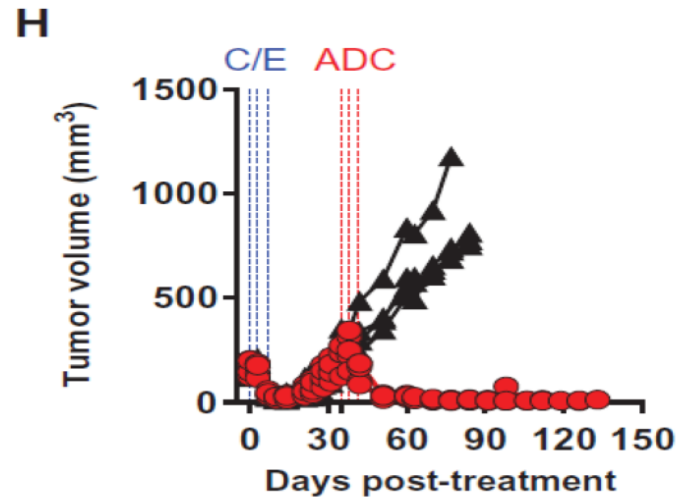


# Phase II result of Rova-T

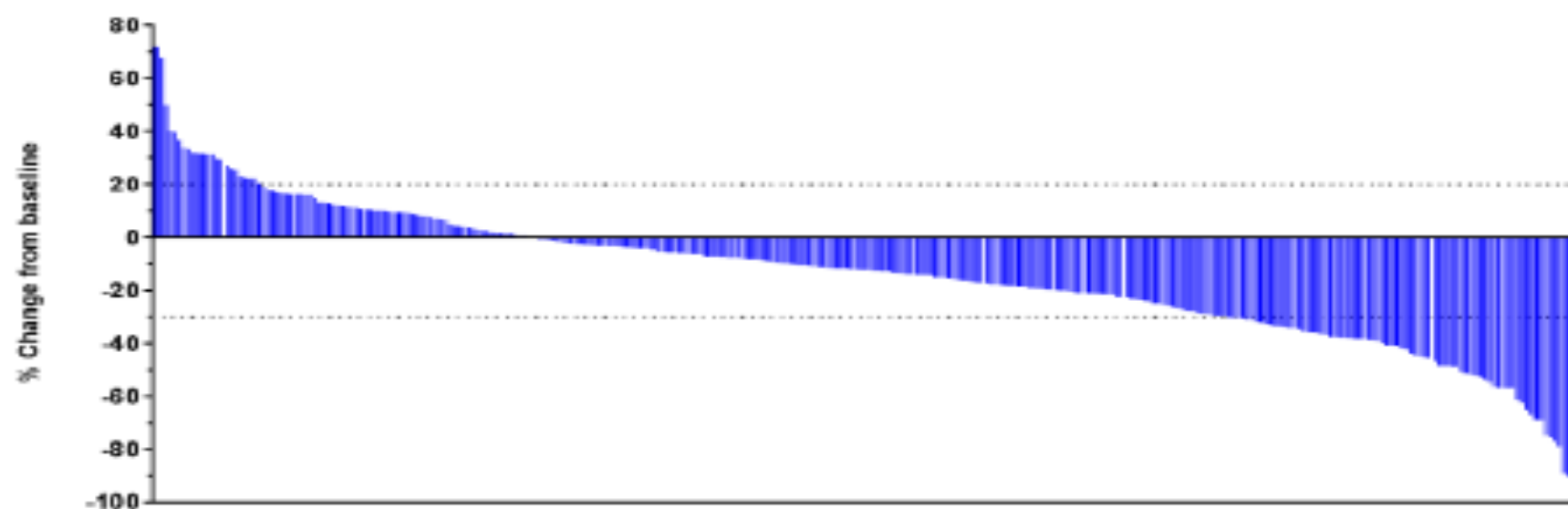
Rova-T: a DLL3 targeting antibody-drug conjugate



rovalpituzumab tesirine (*Rova-T*)



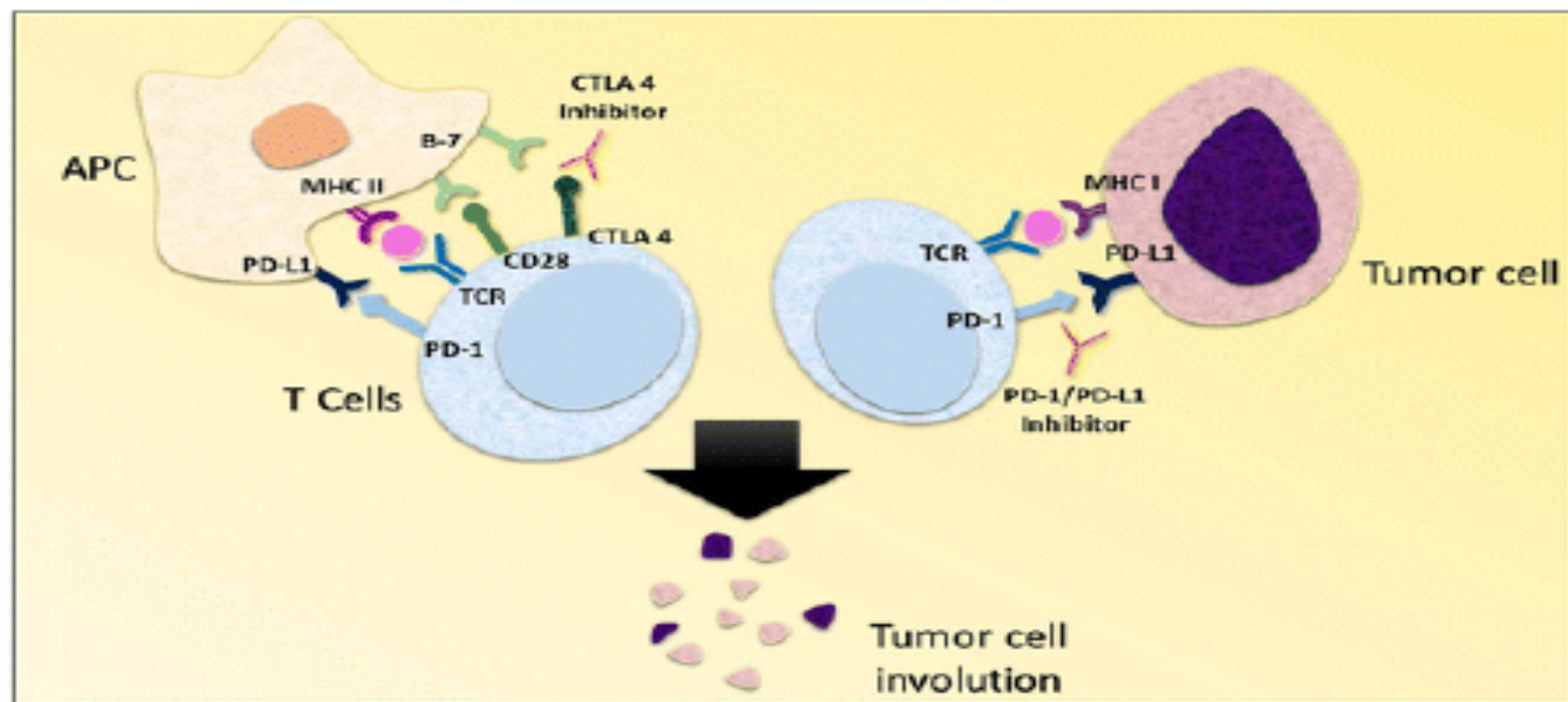
## Phase II result of Rova-T (TRINITY Trial)



ORRs were 12.4%, 14.3% and 13.2% in all, DLL3<sup>high</sup>, and DLL3<sup>+</sup> patients, respectively. Median OS was 5.6 months in all patients.

# Immunotherapy in SCLC

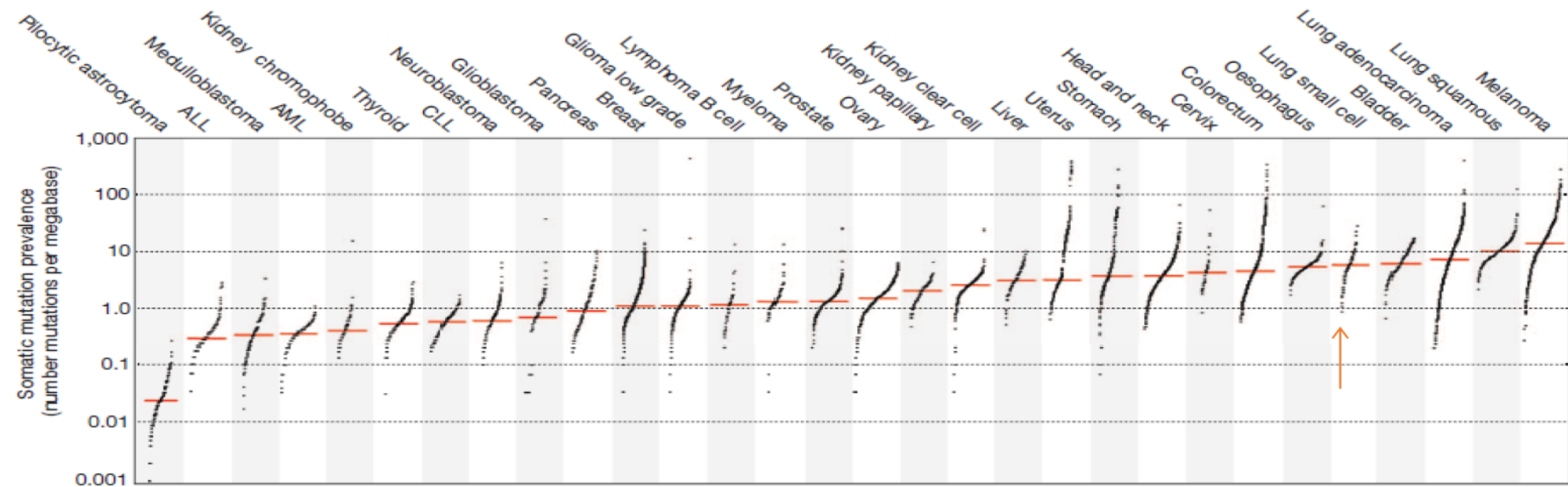
## Immune checkpoints





# Mutation loads

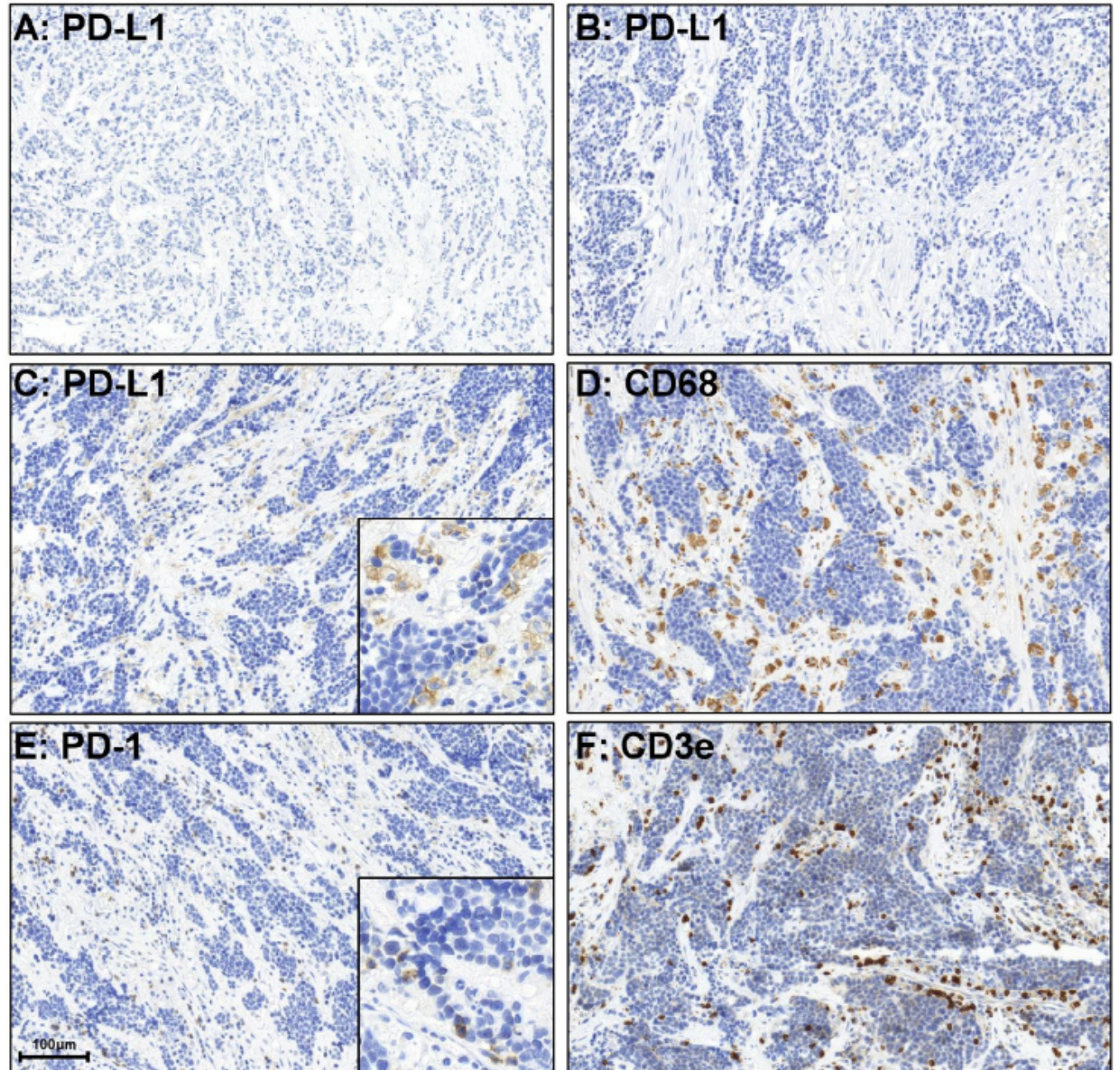
## Mutation loads in different cancer types



**Figure 1 | The prevalence of somatic mutations across human cancer types.** Every dot represents a sample whereas the red horizontal lines are the median numbers of mutations in the respective cancer types. The vertical axis (log scaled) shows the number of mutations per megabase whereas the different

cancer types are ordered on the horizontal axis based on their median numbers of somatic mutations. We thank G. Getz and colleagues for the design of this figure<sup>26</sup>. ALL, acute lymphoblastic leukaemia; AML, acute myeloid leukaemia; CLL, chronic lymphocytic leukaemia.

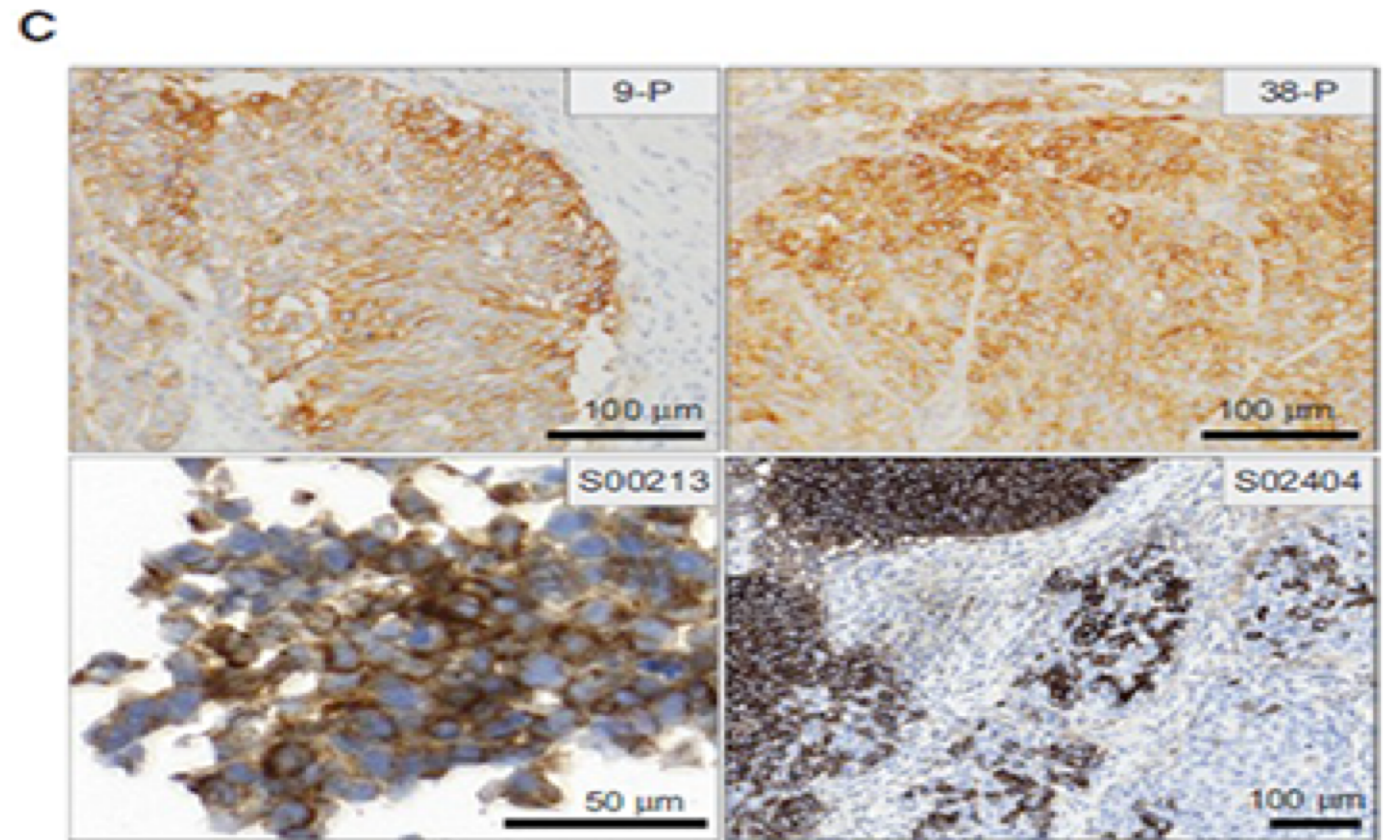
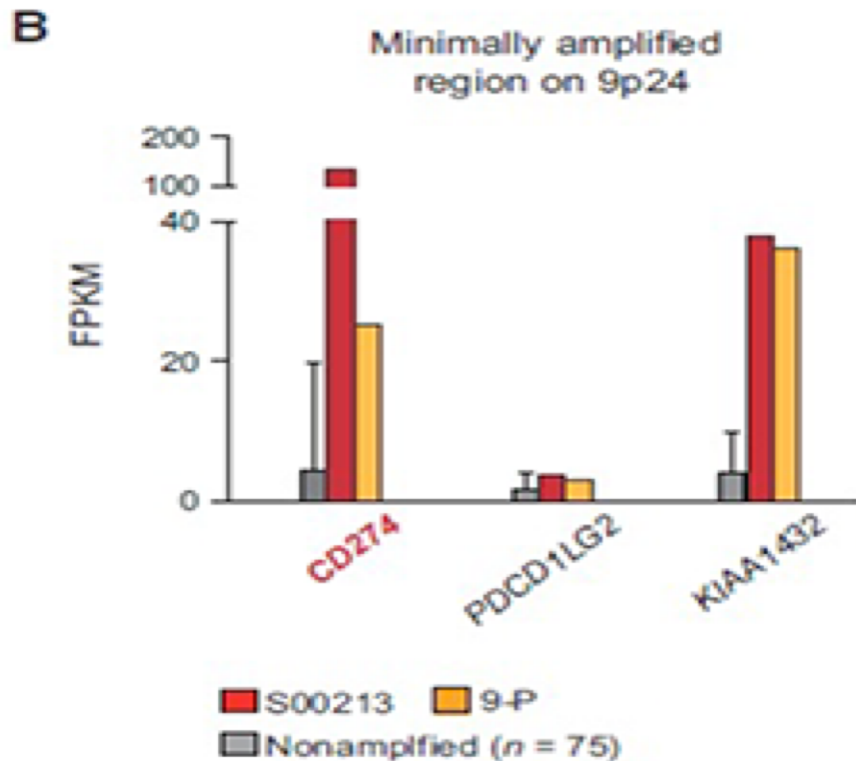
PD-1 and PD-L1  
are expressed in  
the tumor  
stroma of small  
cell carcinoma.





# CD274 amplification

CD274 (PD-L1) gene is amplified in 1.9% of SCLC



# KEYNOTE-028

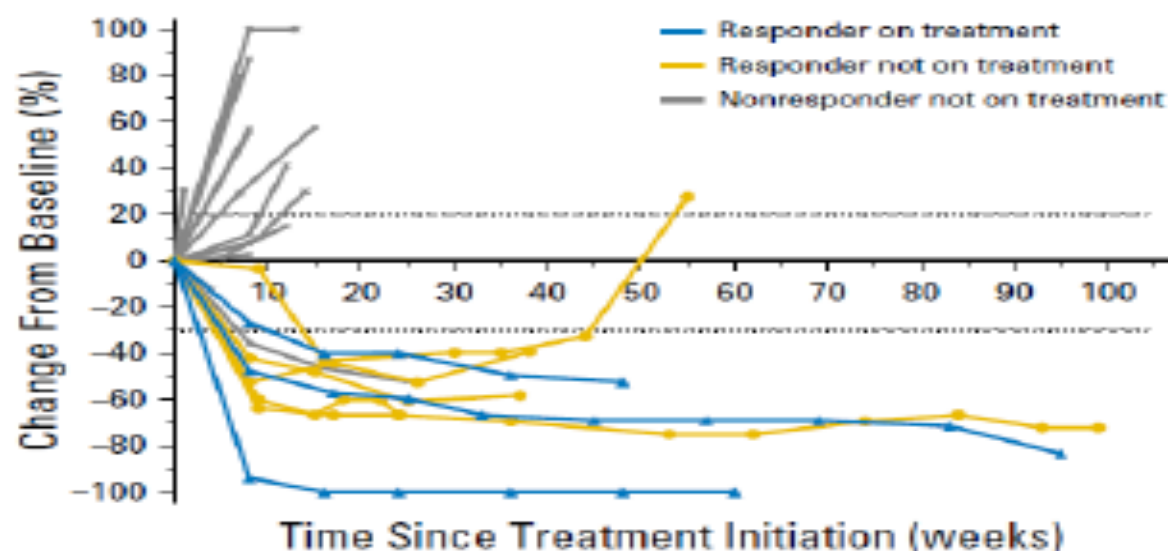
## Third-line: Phase Ib KEYNOTE-028 (Pembrolizumab)

### Study patient population:

1. Histologically confirmed SCLC or pulmonary neuroendocrine tumor
2. PD-L1 expression in  $\geq 1\%$  of tumor and associated inflammatory cells or positive staining in stroma

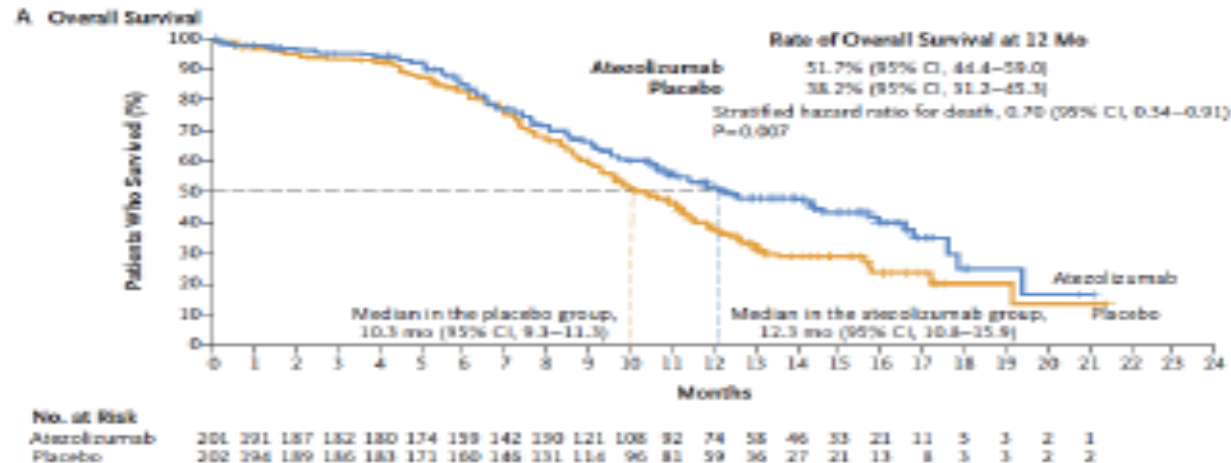
**Table 3.** Confirmed Efficacy Results (investigator-assessed) in the Total Population

Efficacy	Value of Patient Population (n = 24)
ORR*, No. (%) [95% CI]	8 (33.3 [15.6-55.3])
CR, No. (%)	1 (4.2)
PR, No. (%)	7 (29.2)
SD, No. (%)	1 (4.2)
Median DOR, months† (range)	19.4 ( $\geq 3.6$ to $\geq 20.0$ )
Median TTR, months (95% CI)	2.0 (1.7-3.7)
DCR‡, No. (%) [95% CI]	8 (33.3 [15.6-55.3])
Progressive disease, No. (%)	13 (54.2)
Not evaluable, No. (%)	2 (8.3)
PFS	
Events, No. (%)	20 (83.3)
Median, months (95% CI)	1.9 (1.7-5.9)
Six-month rate, % (95% CI)	28.6 (12.4-47.2)
Twelve-month rate, % (95% CI)	23.8 (9.1-42.3)
OS	
Events, No. (%)	15 (62.5)
Median, months (95% CI)	9.7 (4.1-NR)
Six-month rate, % (95% CI)	66.0 (43.3-81.3)
Twelve-month rate, % (95% CI)	37.7 (18.4-57.0)



# Impower 133 trial

## Addition of ICI to front-line chemotherapy improved survival of SCLC patients (Impower 133 trial)



**Table 2. Response Rate, Duration of Response, and Disease Progression.\***

Variable	Atezolizumab Group (N=201)	Placebo Group (N=202)
Objective confirmed response†	121 (60.2 [53.3–67.0])	130 (64.4 [57.3–71.6])
Complete response — no. (%) [95% CI]	5 (2.5 [0.8–5.7])	2 (1.0 [0.3–3.5])
Partial response — no. (%) [95% CI]	116 (57.7 [50.6–64.6])	128 (63.4 [56.3–70.6])
Median duration of response (range) — mo‡	4.2 (1.4–19.5)	3.9 (2.0–16.2)
Ongoing response at data cutoff — no./total no. (%)	18/121 (14.9)	7/130 (5.4)
Stable disease — no. (%) [95% CI]	42 (20.9 [15.5–27.2])	43 (21.3 [15.9–27.6])
Progressive disease — no. (%) [95% CI]	22 (10.9 [7.0–16.1])	14 (6.9 [3.8–11.4])



# Summary

- SCLC is a recalcitrant cancer and new therapy is urgently needed.
- Inactivation of TP53 and RB1 are almost universal in SCLC.
- Newer therapies are on the horizon: Rova-T ADC and Immunotherapy with immune checkpoint inhibitors



Questions?